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Search for 09/530,363

(FILE 'HOME' ENTERED AT 09:22:02 ON 13 DEC 2002)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, BIOTECHDS' ENTERED AT
09:22:26 ON
13 DEC 2002

L1 187 S GABERT, J?/AU
L2 9 S L1 AND (ANCHOR? OR MLL)
L3 6 DUP REM L2 (3 DUPLICATES REMOVED)

L5 2145 S ANCHOR? (10A) (PCR OR AMPLIF? OR POLYMERASE CHAIN OR
PRIMER)
L6 0 S L5 AND BREAKPOINT AND FUSION
L7 146 S L5 AND (TRANSLOCATION OR REARRANGEMENT)
L8 55 DUP REM L7 (91 DUPLICATES REMOVED)

(FILE 'HOME' ENTERED AT 20:27:07 ON 01 JUN 2003)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, BIOTECHDS' ENTERED AT
20:27:24 ON
01 JUN 2003

L1 1240 S ANCHOR AND (TRANSLOCATION OR REARRANGEMENT)
L2 23 S L1 AND (MLL OR BCR)
L3 3309 S ANCHOR? (S) (PRIMER OR PCR OR POLYMERASE CHAIN)
L4 185 S L3 AND (TRANSLOCATION OR REARRANGEMENT)
L5 81 DUP REM L4 (104 DUPLICATES REMOVED)
L6 41 S L3 AND REVIEW
L7 38 DUP REM L6 (3 DUPLICATES REMOVED)

WEST

Freeform Search

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Term:

L6 and 14

 Display: Documents in Display Format: Starting with Number

 Generate: ☐ Hit List ☒ Hit Count ☐ Side by Side ☐ Image

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DB=USPT,PGPB,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ

| | | | |
|-----------|--|------|-----------|
| <u>L8</u> | L6 and 14 | 141 | <u>L8</u> |
| <u>L7</u> | L6 and (breakpoint or fusion or junction or mll) and rearrangement | 1280 | <u>L7</u> |
| <u>L6</u> | specific same (random or anchor\$ or arbitrary or non-specific or degenerate or universal) same primer | 7512 | <u>L6</u> |
| <u>L5</u> | L4 and ((anchor\$ or arbitrary or non-specific or random) with primer) | 172 | <u>L5</u> |
| <u>L4</u> | (breakpoint or fusion or junction or mll) same rearrangement | 1128 | <u>L4</u> |
| <u>L3</u> | L2 and ((arbitrary or non-specific or random) with primer) | 181 | <u>L3</u> |
| <u>L2</u> | L1 and (breakpoint or fusion or junction or mll) and rearrangement | 1544 | <u>L2</u> |
| <u>L1</u> | anchor\$ with primer | 3581 | <u>L1</u> |

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| bcr-abl and (anchor\$ same (primer or pcr or amplific\$ or polymerase chain)) | 21 |

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L5

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| | |
|-----------|---|
| <u>L5</u> | bcr-abl and (anchor\$ same (primer or pcr or amplific\$ or polymerase chain)) |
| <u>L4</u> | bcr-abl and anchor\$ |
| <u>L3</u> | mll gene and anchor\$ |
| <u>L2</u> | mll gene with detect\$ |
| <u>L1</u> | ribozymes MLL gene-specific |

| | |
|-----|-----------|
| 21 | <u>L5</u> |
| 148 | <u>L4</u> |
| 12 | <u>L3</u> |
| 15 | <u>L2</u> |
| 0 | <u>L1</u> |

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| L7 and (primer same target) | 17 |

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| | | | |
|-----------|----------------------------------|------|-----------|
| <u>L8</u> | L7 and (primer same target) | 17 | <u>L8</u> |
| <u>L7</u> | probe same fusion partner | 72 | <u>L7</u> |
| <u>L6</u> | probe same fusion | 3094 | <u>L6</u> |
| <u>L5</u> | L4 same target | 0 | <u>L5</u> |
| <u>L4</u> | anchor\$ primer same fusion gene | 836 | <u>L4</u> |
| <u>L3</u> | L1 same fusion gene | 0 | <u>L3</u> |
| <u>L2</u> | L1 same probe | 14 | <u>L2</u> |
| <u>L1</u> | anchor\$ primer same kit | 308 | <u>L1</u> |

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| L2 same (translocation or rearrangement) | 1 |

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| | | | |
|-----------|---|------|-----------|
| <u>L8</u> | L2 same (translocation or rearrangement) | 1 | <u>L8</u> |
| <u>L7</u> | L2 and mll gene | 1 | <u>L7</u> |
| <u>L6</u> | L2 same (cancer or patholog\$ or carcinoma or mll) | 0 | <u>L6</u> |
| <u>L5</u> | L2 and ((indiscriminat\$ or random or arbitrary) near7 (dna or cdna)) | 78 | <u>L5</u> |
| <u>L4</u> | L2 and ((indiscriminat\$ or random or arbitrary) same (dna or cdna)) | 219 | <u>L4</u> |
| <u>L3</u> | L2 same (indiscriminat\$ or random or arbitrary) | 1 | <u>L3</u> |
| <u>L2</u> | L1 same fusion gene | 840 | <u>L2</u> |
| <u>L1</u> | anchor\$ near6 (primer or pcr or amplific\$ or polymerase chain) | 3009 | <u>L1</u> |

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DB=USPT; PLUR=YES; OP=ADJ

L17 5948624.pn.
L16 L15 and (rearrangement or translocation)
L15 L14 and primer
L14 L6 same l7
L13 5143854.pn.
L12 L11 and breakpoint
L11 L10 and kit
L10 L9 and (probe with biotin)
L9 L8 and (rearrangement or translocation)
L8 l6 and L7
L7 probe with (support or chip or array)
L6 probe same biotin same streptavidin\$
L5 L1 and fusion
L4 L3 and (probe same label)
L3 6309822.pn.
L2 L1 and (rearrangement or translocation)
L1 primers same probe same kit same (support or chip or array)

Hit Count Set Name

result set

1 L17
 40 L16
 128 L15
 151 L14
 1 L13
 11 L12
 204 L11
 226 L10
 252 L9
 678 L8
 15880 L7
 1537 L6
 35 L5
 1 L4
 1 L3
 25 L2
 120 L1

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side by side

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result set

DB=USPT; PLUR=YES; OP=ADJ

| | | | |
|------------|--|------|------------|
| <u>L21</u> | L20 and (probe same (solid support or chip or array)) | 17 | <u>L21</u> |
| <u>L20</u> | L16 and (anchor\$ primer or oligodt or oligo(T)) | 90 | <u>L20</u> |
| <u>L19</u> | L18 and l2 | 75 | <u>L19</u> |
| <u>L18</u> | L17 and (probe same (solid support or chip or array)) | 143 | <u>L18</u> |
| <u>L17</u> | L16 and (anchor\$ primer or oligodt or oligo(T) or oligo\$ primer) | 560 | <u>L17</u> |
| <u>L16</u> | (cancer or leukemia) same (rearrangement or translocation) | 1425 | <u>L16</u> |
| <u>L15</u> | kit same l1 same probe | 12 | <u>L15</u> |
| <u>L14</u> | L13 and (probe same solid support or chip or array) | 23 | <u>L14</u> |
| <u>L13</u> | L12 and l1 | 76 | <u>L13</u> |
| <u>L12</u> | (rearrangement or translocation or bcr-abl or mll) same (cancer or leukemia) | 1592 | <u>L12</u> |
| <u>L11</u> | L4 and dna chip | 11 | <u>L11</u> |
| <u>L10</u> | L8 and (rearrangement or translocation) | 20 | <u>L10</u> |
| <u>L9</u> | L8 and breakpoint | 1 | <u>L9</u> |
| <u>L8</u> | L1 same probe | 55 | <u>L8</u> |
| <u>L7</u> | L6 and l2 | 31 | <u>L7</u> |
| <u>L6</u> | kit same (anchor\$ primer) | 141 | <u>L6</u> |

DB=USPT,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ

| | | | |
|-----------|---|------|-----------|
| <u>L5</u> | L4 and (rearrangement or translocation or breakpoint) | 108 | <u>L5</u> |
| <u>L4</u> | L3 and (solid support or chip or array) | 160 | <u>L4</u> |
| <u>L3</u> | L1 and L2 | 221 | <u>L3</u> |
| <u>L2</u> | biotin\$ with probe | 4994 | <u>L2</u> |
| <u>L1</u> | anchor\$ primer | 786 | <u>L1</u> |

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DB=USPT; PLUR=YES; OP=ADJ

| | |
|------------|--|
| <u>L14</u> | 6309822.pn. and covalent |
| <u>L13</u> | 6040138.pn. or 5800992.pn. or 6309822.pn. |
| <u>L12</u> | L11 and (dna chip or array) |
| <u>L11</u> | fodor.in. |
| <u>L10</u> | 5807522.pn. |
| <u>L9</u> | brown.in. and array |
| <u>L8</u> | L7 and anchor\$ primer |
| <u>L7</u> | (biotin same support) same streptavidin\$ same probe |
| <u>L6</u> | L5 and anchor\$ primer |
| <u>L5</u> | (biotin with support) same streptavidin\$ same probe |
| <u>L4</u> | (biotin same support) same strepavidin\$ same probe |
| <u>L3</u> | (biotin with support) same strepavidin\$ same probe |
| <u>L2</u> | mll and exon 5 |
| <u>L1</u> | mll and rt-pcr |

Hit Count Set Name

result set

| | |
|------|------------|
| 1 | <u>L14</u> |
| 3 | <u>L13</u> |
| 84 | <u>L12</u> |
| 207 | <u>L11</u> |
| 1 | <u>L10</u> |
| 1384 | <u>L9</u> |
| 6 | <u>L8</u> |
| 177 | <u>L7</u> |
| 6 | <u>L6</u> |
| 100 | <u>L5</u> |
| 5 | <u>L4</u> |
| 4 | <u>L3</u> |
| 13 | <u>L2</u> |
| 36 | <u>L1</u> |

END OF SEARCH HISTORY

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Search for 09/530,363

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L3 6 DUP REM L2 (3 DUPLICATES REMOVED)

L5 2145 S ANCHOR? (10A) (PCR OR AMPLIF? OR POLYMERASE CHAIN OR
PRIMER)

L6 0 S L5 AND BREAKPOINT AND FUSION

L7 146 S L5 AND (TRANSLOCATION OR REARRANGEMENT)

L8 55 DUP REM L7 (91 DUPLICATES REMOVED)

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side by side

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result set

DB=USPT,PGPB,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ

| | | | |
|------------|--|------|------------|
| <u>L17</u> | L14 same (translocation or rearrangement) | 15 | <u>L17</u> |
| <u>L16</u> | L14 and (translocation or rearrangement) | 1193 | <u>L16</u> |
| <u>L15</u> | L14 and (breakpoint same fusion) | 7 | <u>L15</u> |
| <u>L14</u> | (reverse anchor\$ or anchor\$) same (pcr or amplification or polymerase chain) | 2491 | <u>L14</u> |
| <u>L13</u> | (reverse anchor\$ or anchor) same (pcr or amplification or polymerase chain) | 1620 | <u>L13</u> |
| <u>L12</u> | L11 and anchor\$ | 8 | <u>L12</u> |
| <u>L11</u> | L10 and (translocation or rearrangement) | 25 | <u>L11</u> |
| <u>L10</u> | mll gene | 27 | <u>L10</u> |

DB=USPT; PLUR=YES; OP=ADJ

| | | | |
|-----------|---|------|-----------|
| <u>L9</u> | L8 and (breakpoint same fusion) | 4 | <u>L9</u> |
| <u>L8</u> | (anchor\$ same primer) and (translocation or rearrangement) | 485 | <u>L8</u> |
| <u>L7</u> | L4 and (breakpoint same fusion) | 3 | <u>L7</u> |
| <u>L6</u> | L4 and myeloid lymphoid leukemia | 0 | <u>L6</u> |
| <u>L5</u> | L4 and mll gene | 0 | <u>L5</u> |
| <u>L4</u> | L3 and (translocation or rearrangement) | 509 | <u>L4</u> |
| <u>L3</u> | anchor\$ same (pcr or polymerase chain) | 1289 | <u>L3</u> |
| <u>L2</u> | (anchor\$ (pcr or polymerase chain or amplif\$)).ab. | 1 | <u>L2</u> |
| <u>L1</u> | (anchor\$ (pcr or polymerase chain or amplif\$)).ti. | 0 | <u>L1</u> |

END OF SEARCH HISTORY



| | | | | | | | | |
|---------------|-------------------------------------|---------------|--------|-----------|--------|-----------|------|---------|
| PubMed | Nucleotide | Protein | Genome | Structure | PopSet | Taxonomy | OMIM | Books |
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- Search numbers may not be continuous; all searches are represented.

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| #13 | Search anchored pcr and review and mll | 11:22:53 | <u>6</u> |
| #11 | Related Articles for PubMed (Select 9372194) | 11:21:19 | <u>150</u> |
| #9 | Search anchor* primer AND review | 11:19:34 | <u>5</u> |
| #7 | Search anchored primer and review | 11:18:46 | <u>1</u> |
| #3 | Search anchored pcr AND review AND rearrangement* and primer | 11:16:34 | <u>2</u> |
| #2 | Search anchored pcr AND review | 11:15:55 | <u>4410</u> |
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